If you

- know the structure
- have observed all of the variables
- have no missing data
- you can learn each conditional probability separately.
- \longrightarrow supervised learning



Learning conditional probabilities

- Each conditional probability distribution can be learned separately:
- For example:

$$P(E = t \mid A = t \land B = f)$$

=
$$\frac{(\#\text{examples: } E = t \land A = t \land B = f) + c_1}{(\#\text{examples: } A = t \land B = f) + c}$$

where c_1 and c reflect prior (expert) knowledge ($c_1 \leq c$).

• When there are many parents to a node, there can little or no data for each conditional probability: use supervised learning to learn a decision tree, linear classifier, a neural network or other representation of the conditional probability.

Unobserved Variables



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- Repeat the following two steps:
 - E-step give the expected number of data points for the unobserved variables based on the given probability distribution. Requires probabilistic inference.
 - M-step infer the (maximum likelihood) probabilities from the data. This is the same as the fully-observable case.
- Start either with made-up data or made-up probabilities.
- EM will converge to a local maxima.

Belief network structure learning (I)

Given examples **e**, and model *m*:

$$P(m \mid \mathbf{e}) = rac{P(\mathbf{e} \mid m) * P(m)}{P(\mathbf{e}).}$$

- A model here is a belief network.
- A bigger network can always fit the data better.
- *P*(*m*) lets us encode a preference for simpler models (e.g, smaller networks)
- \longrightarrow search over network structure looking for the most likely model.
 - Taking logarithms (and negating):

$$\arg \max_{m} P(m \mid \mathbf{e}) = \arg \min_{m} (-\log P(\mathbf{e} \mid m) - \log P(m))$$

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Bayes Rule:

 $P(h|d) \propto P(d|h)P(h)$ arg max $P(h|d) = \arg \max_{h} P(d|h)P(h)$ $= \arg \max_{h} (\log P(d|h) + \log P(h))$

- $\log P(d|h)$ measures fit to data
- $\log P(h)$ measures model complexity

- A bit is a binary digit.
- 1 bit can distinguish 2 items
- k bits can distinguish 2^k items
- n items can be distinguished using $\log_2 n$ bits
- Can you do better?

Consider a code to distinguish elements of $\{a, b, c, d\}$ with

$$P(a) = \frac{1}{2}, P(b) = \frac{1}{4}, P(c) = \frac{1}{8}, P(d) = \frac{1}{8}$$

Consider the code:

This code uses 1 to 3 bits. On average, it uses

$$P(a) * 1 + P(b) * 2 + P(c) * 3 + P(d) * 3$$

= $\frac{1}{2} + \frac{2}{4} + \frac{3}{8} + \frac{3}{8} = 1\frac{3}{4}$ bits.

The string *aacabbda* has code 00110010101110. The code 0111110010100 represents string *adcabba*

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Information Content

- To identify x, you need $-\log_2 P(x)$ bits.
- Given a distribution over a set of values, to identify a member, the expected number of bits is

$$\sum_{x} -P(x) * \log_2 P(x)$$

is the information content or entropy of the distribution.

• The expected number of bits it takes to describe a distribution given evidence *e*:

$$I(e) = \sum_{x} -P(x|e) * \log_2 P(x|e).$$

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Given a test that can distinguish the cases where α is true from the cases where α is false, the information gain from this test is:

$$I(true) - (P(\alpha) * I(\alpha) + P(\neg \alpha) * I(\neg \alpha)).$$

- *I*(*true*) is the expected number of bits needed before the test
- P(α) * I(α) + P(¬α) * I(¬α) is the expected number of bits after the test.

Comparing Distributions

- Suppose a code is designed to be optimal for probability distribution Q, so that x is described using -log₂Q(x) bits.
- Suppose *P* is another probability distribution. The expected number of bits to describe *P* using the code for *Q* is

$$\sum_{x} -P(x) * \log_2 Q(x)$$

 The difference between this and the entropy of P describing P using its optimal code – is the Kullback–Leibler (KL) divergence (also called relative entropy):

$$D_{KL}(P \mid\mid Q) = (\sum_{x} -P(x) * \log Q(x)) - \sum_{x} -P(x) * \log P(x)$$
$$= \sum_{x} -P(x) * \log(Q(x)/P(x))$$

• When is this large? When $P(x) \gg Q(x)$ for some x.

- Search over total orderings of variables.
- For each total ordering X₁,..., X_n use supervised learning to learn P(X_i | X₁...X_{i-1}).
- Return the network model found with minimum:
 - $-\log P(\mathbf{e} \mid m) \log P(m)$
 - $P(\mathbf{e} \mid m)$ can be obtained by inference.
 - How to determine $-\log P(m)$?

Bayesian Information Criterion (BIC) Score

$$P(m \mid \mathbf{e}) = \frac{P(\mathbf{e} \mid m) * P(m)}{P(\mathbf{e})}$$
$$-\log P(m \mid \mathbf{e}) \propto -\log P(\mathbf{e} \mid m) - \log P(m)$$

- -log P(e | m) is the negative log likelihood of model m: number of bits to describe the data in terms of the model.
- $|\mathbf{e}|$ is the number of examples. Each proposition can be true for between 0 and $|\mathbf{e}|$ examples, so there are $|\mathbf{e}| + 1$ different probabilities to distinguish. Each one can be described in $\log(|\mathbf{e}| + 1) \approx \log(|\mathbf{e}|)$ bits.
- If there are ||m|| independent parameters (||m|| is the dimensionality of the model):

 $-\log P(m \mid \mathbf{e}) \propto -\log P(\mathbf{e} \mid m) + ||m||\log(|\mathbf{e}|)$

This is the Bayesian Information Criterion (BIC) score.

- Given a total ordering, to determine *parents*(X_i) do independence tests to determine which features should be the parents
- XOR problem: just because features do not give information individually, does not mean they will not give information in combination
- Search over total orderings of variables

- You cannot just ignore missing data unless you know it is missing at random.
- Is the reason data is missing correlated with something of interest?
- For example: data in a clinical trial to test a drug may be missing because:
 - the patient dies
 - the patient had severe side effects
 - the patient was cured
 - the patient had to visit a sick relative.

— ignoring some of these may make the drug look better or worse than it is.

• In general you need to model why data is missing (see Chapter 11)

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- We have a mixture of observational data and data from randomized studies.
- We are not given the structure.
- We don't know whether there are hidden variables or not. We don't know the domain size of hidden variables.
- There is missing data.
- ... this is too difficult for current techniques!